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piece 1, NC_000913, yhjA_treF+, config: linear, direction: +, begin: 3667182, end: 3667634

The figure shows a genomic map of the *yhjA*-*treF+* locus. The top line represents the DNA sequence with transcription start sites marked by asterisks (*). The bottom line shows the deduced amino acid sequence for each gene product. Red dots indicate specific mutations or features. The *yhjA* gene encodes a protein with a putative signal peptide (red box) and a trehalose-6-phosphate hydrolase domain (green box). The *treF+* gene encodes a protein with a putative signal peptide (red box) and a trehalose-6-phosphate hydrolase domain (green box).

p35 1.4 bits

{-----} sd-(10)-ir 3667240 Gap 2.7 bits

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|-----| sd-ir 3667240 yhjA_treF+ total 7.3 bits  
|-----| p35-(23)-p10 3667237 Gap 4.4 bits  
|-----| p35-p10 3667237 total 4.5 bits
```

sd ir yhjA_treF+ [### ... or

|-----| sd-ir 3667319 yhjA_treF+ total 7.7 bits ... p10

The diagram illustrates the p35 gene structure. It features a dashed black line at the top representing the genomic DNA. Below it, a solid blue line represents the transcribed mRNA. The gene is divided into several exons, shown as colored boxes: a small red box, a large green box, a smaller green box, a yellow box, a purple box, and a pink box. Introns are represented by grey boxes between the exons. A red bracket labeled 'sd' indicates the start site of transcription. A red arrow points to a specific position within the first exon, which is highlighted with a red box. To the right of the gene structure, the text 'p35 2.4 bit' is displayed.

{-----} sd-(15)-ir 3667341 Gap 6.0 bits
----- sd-ir 3667341 yhja_treF+ total 8.5
 p35 5.6 bits {-----} ... p35-(23)-p10 3667360 Gap

... p35-(24)-p10 3667344 Gap
... p35-p10 3667344 total 5.4

0x0000000000000000 p35 1.3 bits |----- ... p35-p10 3667360 total 6.7
0x0000000000000000 {----- ... p35-(24)-p10 3667349 Gap

... p35-p10 3667349 total 5.4
... p35

... > orf 35 codons



